

# SCORE Search Results Details for Application 10560250 and Search Result 20071114\_085447\_us-10-560-250-48.rapbm.

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This page gives you Search Results detail for the Application 10560250 and Search Result 20071114\_085447\_us-10-560-250-48.rapbm.

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GenCore version 6.2.1  
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OM protein - protein search, using sw model

Run on: November 14, 2007, 08:54:24 ; Search time 133 Seconds  
(without alignments)  
98.389 Million cell updates/sec

Title: US-10-560-250-48  
Perfect score: 82  
Sequence: 1 IGINVRAREGLTFDND 16

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 3552611 seqs, 817857308 residues

Total number of hits satisfying chosen parameters: 3552611

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Published\_Applications\_AA\_Main:\*

- 1: /EMC\_Celerra\_SIDS2/ptodata/1/pubpaa/US07\_PUBCOMB.pep:\*
- 2: /EMC\_Celerra\_SIDS2/ptodata/1/pubpaa/US08\_PUBCOMB.pep:\*
- 3: /EMC\_Celerra\_SIDS2/ptodata/1/pubpaa/US09\_PUBCOMB.pep:\*
- 4: /EMC\_Celerra\_SIDS2/ptodata/1/pubpaa/US10A\_PUBCOMB.pep:\*
- 5: /EMC\_Celerra\_SIDS2/ptodata/1/pubpaa/US10B\_PUBCOMB.pep:\*
- 6: /EMC\_Celerra\_SIDS2/ptodata/1/pubpaa/US11\_PUBCOMB.pep:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result	Query	%				
No.	Score	Match Length	DB	ID		Description

1	82	100.0	16	5	US-10-560-250-48	Sequence 48, Appl
2	82	100.0	365	5	US-10-560-250-37	Sequence 37, Appl
3	82	100.0	365	5	US-10-560-250-39	Sequence 39, Appl
4	82	100.0	366	5	US-10-560-250-57	Sequence 57, Appl
5	82	100.0	380	4	US-10-040-949A-47	Sequence 47, Appl
6	82	100.0	381	3	US-09-348-354A-31	Sequence 31, Appl
7	82	100.0	381	3	US-09-953-280-31	Sequence 31, Appl
8	82	100.0	381	6	US-11-207-626A-31	Sequence 31, Appl
9	82	100.0	578	5	US-10-560-250-55	Sequence 55, Appl